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10/519328

SEQUENCE LISTING

<110> BOUGUELERET; Lydie
NIKNEJAD; Anne

<120> SECRETED PEPTIDES

<130> 4-33619A/GEP

<140> 10/519,328

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<150> PCT/EP03/007069

<151> 2003-07-02

<150> 60/393,840

<151> 2002-07-02

<160> 10

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 1821

<212> PRT

<213> Homo sapiens

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<221> PROPEP

<222> (1)...(1821)

<223> Latent transforming growth factor-beta-binding
protein precursor 2

<221> SIGNAL

<222> (1)...(35)

<223> Predicted by SignalP version 2.0

<221> SITE

<222> (96)...(97)

<223> Dibasic peptidase cleavage site

<221> SITE

<222> (104)...(105)

<223> Dibasic peptidase cleavage site

<221> SITE

<222> (113)...(114)

<223> Dibasic peptidase cleavage site

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Gly	Arg	Leu	Thr	Gly	Arg	Asn	Val	Cys	Gly	Gln	Cys	Cys	Pro	Gly	
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Glu	Arg	Ser	Pro	Asn	Leu	Arg	Arg	Ser	Ser	Ala	Ala	Gly	Glu	Gly	Thr
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Gly	Ser	Tyr	Arg	Cys	Val	Leu	Gly	Cys	Gln	Pro	Gly	Phe	His	Met	Ala
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Val	Tyr	Ser	Leu	Phe	Arg	Glu	Gln	Asp	Ala	Pro	Val	Ala	Gly	Leu	Gln		
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Pro	Val	Glu	Arg	Ala	Gln	Pro	Gly	Trp	Gly	Ser	Pro	Arg	Arg	Pro	Thr		
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Gly Phe Arg Gly Ala Arg Cys Glu Glu Val Ile Pro Asp Glu Glu Phe		175
	180	185
Asp Pro Gln Asn Ser Arg Leu Ala Pro Arg Arg Trp Ala Glu Arg Ser		190
	195	200
Pro Asn Leu Arg Arg Ser Ser Ala Ala Gly Glu Gly Thr Leu Ala Arg		205
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Ala Gln Pro Pro Ala Pro Gln Ser Pro Pro Ala Pro Gln Ser Pro Pro		220
	225	230
Ala Gly Thr Leu Ser Gly Leu Ser Gln Thr His Pro Ser Gln Gln His		235
	245	250
Val Gly Leu Ser Arg Thr Val Arg Leu His Pro Thr Ala Thr Ala Ser		255
	260	265
Ser Gln Leu Ser Ser Asn Ala Leu Pro Pro Gly Pro Gly Leu Glu Gln		270
	275	280
Arg Asp Gly Thr Gln Gln Ala Val Pro Leu Glu His Pro Ser Ser Pro		285
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Trp Gly Leu Asn Leu Thr Glu Lys Ile Lys Lys Ile Lys Ile Val Phe		300
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Thr Pro Thr Ile Cys Lys Gln Thr Cys Ala Arg Gly His Cys Ala Asn		315
	325	330
Ser Cys Glu Arg Gly Asp Thr Thr Thr Leu Tyr Ser Gln Gly Gly His		335
	340	345
Gly His Asp Pro Lys Ser Gly Phe Arg Ile Tyr Phe Cys Gln Ile Pro		350
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Cys Leu Asn Gly Gly Arg Cys Ile Gly Arg Asp Glu Cys Trp Cys Pro		365
	370	375
Ala Asn Ser Thr Gly Lys Phe Cys His Leu Pro Ile Pro Gln Pro Asp		380
	385	390
Arg Glu Pro Pro Gly Arg Gly Ser Arg Pro Arg Ala Leu Leu Glu Ala		395
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Pro Leu Lys Gln Ser Thr Phe Thr Leu Pro Leu Ser Asn Gln Leu Ala		415
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Ser Val Asn Pro Ser Leu Val Lys Val His Ile His His Pro Pro Glu		430
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Ala Ser Val Gln Ile His Gln Val Ala Gln Val Arg Gly Gly Val Glu		445
	450	455
Glu Ala Leu Val Glu Asn Ser Val Glu Thr Arg Pro Pro Pro Trp Leu		460
	465	470
Pro Ala Ser Pro Gly His Ser Leu Trp Asp Ser Asn Asn Ile Pro Ala		475
	485	490
Arg Ser Gly Glu Pro Pro Arg Pro Leu Pro Pro Ala Ala Pro Arg Pro		495
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Arg Gly Leu Leu Gly Arg Cys Tyr Leu Asn Thr Val Asn Gly Gln Cys		510
	515	520
Ala Asn Pro Leu Leu Glu Leu Thr Thr Gln Glu Asp Cys Cys Gly Ser		525
	530	535
Val Gly Ala Phe Trp Gly Val Thr Leu Cys Ala Pro Cys Pro Pro Arg		540
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Pro Ala Ser Pro Val Ile Glu Asn Gly Gln Leu Glu Cys Pro Gln Gly		555
	565	570
Tyr Lys Arg Leu Asn Leu Thr His Cys Gln Asp Ile Asn Glu Cys Leu		575
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Thr Leu Gly Leu Cys Lys Asp Ala Glu Cys Val Asn Thr Arg Gly Ser		590
	595	600
Tyr Leu Cys Thr Cys Arg Pro Gly Leu Met Leu Asp Pro Ser Arg Ser		605
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		620

Arg	Cys	Val	Ser	Asp	Lys	Ala	Ile	Ser	Met	Leu	Gln	Gly	Leu	Cys	Tyr	625	630	635	640
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Glu	Cys	Glu	Lys	Cys	Pro	Leu	Pro	Gly	Thr	Glu	Ala	Phe	Arg	Glu	Ile	675	680	685	
Cys	Pro	Ala	Gly	His	Gly	Tyr	Thr	Tyr	Ala	Ser	Ser	Asp	Ile	Arg	Leu	690	695	700	
Ser	Met	Arg	Lys	Ala	Glu	Glu	Glu	Glu	Leu	Ala	Arg	Pro	Pro	Arg	Glu	705	710	715	720
Gln	Gly	Gln	Arg	Ser	Ser	Gly	Ala	Leu	Pro	Gly	Pro	Ala	Glu	Arg	Gln	725	730	735	
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Asp	Lys	Gly	Asp	Ser	Gln	Ala	Gly	Gln	Val	Thr	Thr	Ser	Val	Thr	His	755	760	765	
Ala	Pro	Ala	Trp	Val	Thr	Gly	Asn	Ala	Thr	Thr	Pro	Pro	Met	Pro	Glu	770	775	780	
Gln	Gly	Ile	Ala	Glu	Ile	Gln	Glu	Glu	Gln	Val	Thr	Pro	Ser	Thr	Asp	785	790	795	800
Val	Leu	Val	Thr	Leu	Ser	Thr	Pro	Gly	Ile	Asp	Arg	Cys	Ala	Ala	Gly	805	810	815	
Ala	Thr	Asn	Val	Cys	Gly	Pro	Gly	Thr	Cys	Val	Asn	Leu	Pro	Asp	Gly	820	825	830	
Tyr	Arg	Cys	Val	Cys	Ser	Pro	Gly	Tyr	Gln	Leu	His	Pro	Ser	Gln	Ala	835	840	845	
Tyr	Cys	Thr	Asp	Asp	Asn	Glu	Cys	Leu	Arg	Asp	Pro	Cys	Lys	Gly	Lys	850	855	860	
Gly	Arg	Cys	Ile	Asn	Arg	Val	Gly	Ser	Tyr	Ser	Cys	Phe	Cys	Tyr	Pro	865	870	875	880
Gly	Tyr	Thr	Leu	Ala	Thr	Ser	Gly	Ala	Thr	Gln	Glu	Cys	Gln	Asp	Ile	885	890	895	
Asn	Glu	Cys	Glu	Gln	Pro	Gly	Val	Cys	Ser	Gly	Gly	Gln	Cys	Thr	Asn	900	905	910	
Thr	Glu	Gly	Ser	Tyr	His	Cys	Glu	Cys	Asp	Gln	Gly	Tyr	Ile	Met	Val	915	920	925	
Arg	Lys	Gly	His	Cys	Gln	Asp	Ile	Asn	Glu	Cys	Arg	His	Pro	Gly	Thr	930	935	940	
Cys	Pro	Asp	Gly	Arg	Cys	Val	Asn	Ser	Pro	Gly	Ser	Tyr	Thr	Cys	Leu	945	950	955	960
Ala	Cys	Glu	Glu	Gly	Tyr	Arg	Gly	Gln	Ser	Gly	Ser	Cys	Val	Asp	Val	965	970	975	
Asn	Glu	Cys	Leu	Thr	Pro	Gly	Val	Cys	Ala	His	Gly	Lys	Cys	Thr	Asn	980	985	990	
Leu	Glu	Gly	Ser	Phe	Arg	Cys	Ser	Cys	Glu	Gln	Gly	Tyr	Glu	Val	Thr	995	1000	1005	
Ser	Asp	Glu	Lys	Gly	Cys	Gln	Asp	Val	Asp	Glu	Cys	Ala	Ser	Arg	Ala	1010	1015	1020	
Ser	Cys	Pro	Thr	Gly	Leu	Cys	Leu	Asn	Thr	Glu	Gly	Ser	Phe	Ala	Cys	1025	1030	1035	1040
Ser	Ala	Cys	Glu	Asn	Gly	Tyr	Trp	Val	Asn	Glu	Asp	Gly	Thr	Ala	Cys	1045	1050	1055	
Glu	Asp	Leu	Asp	Glu	Cys	Ala	Phe	Pro	Gly	Val	Cys	Pro	Ser	Gly	Val	1060	1065	1070	
Cys	Thr	Asn	Thr	Ala	Gly	Ser	Phe	Ser	Cys	Lys	Asp	Cys	Asp	Gly	Gly	1075	1080	1085	
Tyr	Arg	Pro	Ser	Pro	Leu	Gly	Asp	Ser	Cys	Glu	Asp	Val	Asp	Glu	Cys	1090	1095	1100	
Glu	Asp	Pro	Gln	Ser	Ser	Cys	Leu	Gly	Gly	Glu	Cys	Lys	Asn	Thr	Val				

1105		1110		1115		1120
Gly Ser Tyr Gln Cys	Leu Cys Pro Gln Gly	Phe Gln Leu Ala Asn Gly				
	1125		1130			1135
Thr Val Cys Glu Asp	Val Asn Glu Cys Met Gly	Glu Glu His Cys Ala				
	1140		1145			1150
Pro His Gly Glu Cys	Leu Asn Ser His Gly	Ser Phe Phe Cys Leu Cys				
	1155		1160			1165
Ala Pro Gly Phe Val	Ser Ala Glu Gly Gly	Thr Ser Cys Gln Asp Val				
	1170		1175			1180
Asp Glu Cys Ala Thr	Thr Asp Pro Cys Val Gly	Gly Gly His Cys Val Asn				
	1185		1190			1195
Thr Glu Gly Ser Phe	Asn Cys Leu Cys Glu Thr	Gly Phe Gln Pro Ser				
	1205		1210			1215
Pro Glu Ser Gly Glu	Cys Val Asp Ile Asp	Glu Cys Glu Asp Tyr Gly				
	1220		1225			1230
Asp Pro Val Cys Gly	Thr Trp Lys Cys Glu Asn	Ser Pro Gly Ser Tyr				
	1235		1240			1245
Arg Cys Val Leu Gly	Cys Gln Pro Gly Phe	His Met Ala Pro Asn Gly				
	1250		1255			1260
Asp Cys Ile Asp Ile	Asp Glu Cys Ala Asn	Asp Thr Met Cys Gly Ser				
	1265		1270			1275
His Gly Phe Cys Asp	Asn Thr Asp Gly Ser	Phe Arg Cys Leu Cys Asp				
	1285		1290			1295
Gln Gly Phe Glu Ile	Ser Pro Ser Gly Trp	Asp Cys Val Asp Val Asn				
	1300		1305			1310
Glu Cys Glu Leu Met	Leu Ala Val Cys Gly	Ala Ala Leu Cys Glu Asn				
	1315		1320			1325
Val Glu Gly Ser Phe	Leu Cys Leu Cys Ala	Ser Asp Leu Glu Glu Tyr				
	1330		1335			1340
Asp Ala Gln Glu Gly	His Cys Arg Pro Arg	Gly Ala Gly Gly Gln Ser				
	1345		1350			1355
Met Ser Glu Ala Pro	Thr Gly Asp His Ala	Pro Ala Pro Thr Arg Met				
	1365		1370			1375
Asp Cys Tyr Ser Gly	Gln Lys Gly His Ala	Pro Cys Ser Ser Val Leu				
	1380		1385			1390
Gly Arg Asn Thr Thr	Gln Ala Glu Cys Cys	Cys Thr Gln Gly Ala Ser				
	1395		1400			1405
Trp Gly Asp Ala Cys	Asp Leu Cys Pro Ser	Glu Asp Ser Ala Glu Phe				
	1410		1415			1420
Ser Glu Ile Cys Pro	Ser Gly Lys Gly Tyr	Ile Pro Val Glu Gly Ala				
	1425		1430			1435
Trp Thr Phe Gly Gln	Thr Met Tyr Thr	Asp Ala Asp Glu Cys Val Ile				
	1445		1450			1455
Phe Gly Pro Gly Leu	Cys Pro Asn Gly Arg	Cys Leu Asn Thr Val Pro				
	1460		1465			1470
Gly Tyr Val Cys Leu	Cys Asn Pro Gly Phe	His Tyr Asp Ala Ser His				
	1475		1480			1485
Lys Lys Cys Glu Asp	His Asp Glu Cys Gln	Asp Leu Ala Cys Glu Asn				
	1490		1495			1500
Gly Glu Cys Val Asn	Thr Glu Gly Ser Phe	His Cys Phe Cys Ser Pro				
	1505		1510			1515
Pro Leu Thr Leu Asp	Leu Ser Gln Gln Arg	Cys Met Asn Ser Thr Ser				
	1525		1530			1535
Ser Thr Glu Asp Leu	Pro Asp His Asp	Ile His Met Asp Ile Cys Trp				
	1540		1545			1550
Lys Lys Val Thr Asn	Asp Val Cys Ser Glu	Pro Leu Arg Gly His Arg				
	1555		1560			1565
Thr Thr Tyr Thr Glu	Cys Cys Cys Gln Asp	Gly Glu Ala Trp Ser Gln				
	1570		1575			1580
Gln Cys Ala Leu Cys	Pro Pro Arg Ser Ser	Glu Val Tyr Ala Gln Leu				
	1585		1590			1595
						1600

Cys Asn Val Ala Arg Ile Glu Ala Glu Arg Glu Ala Gly Val His Phe
 1605 1610 1615
 Arg Pro Gly Tyr Glu Tyr Gly Pro Gly Pro Asp Asp Leu His Tyr Ser
 1620 1625 1630
 Ile Tyr Gly Pro Asp Gly Ala Pro Phe Tyr Asn Tyr Leu Gly Pro Glu
 1635 1640 1645
 Asp Thr Val Pro Glu Pro Ala Phe Pro Asn Thr Ala Gly His Ser Ala
 1650 1655 1660
 Asp Arg Thr Pro Ile Leu Glu Ser Pro Leu Gln Pro Ser Glu Leu Gln
 1665 1670 1675 1680
 Pro His Tyr Val Ala Ser His Pro Glu Pro Pro Ala Gly Phe Glu Gly
 1685 1690 1695
 Leu Gln Ala Glu Glu Cys Gly Ile Leu Asn Gly Cys Glu Asn Gly Arg
 1700 1705 1710
 Cys Val Arg Val Arg Glu Gly Tyr Thr Cys Asp Cys Phe Glu Gly Phe
 1715 1720 1725
 Gln Leu Asp Ala Ala His Met Ala Cys Val Asp Val Asn Glu Cys Asp
 1730 1735 1740
 Asp Leu Asn Gly Pro Ala Val Leu Cys Val His Gly Tyr Cys Glu Asn
 1745 1750 1755 1760
 Thr Glu Gly Ser Tyr Arg Cys His Cys Ser Pro Gly Tyr Val Ala Glu
 1765 1770 1775
 Ala Gly Pro Pro His Cys Thr Ala Lys Glu
 1780 1785

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 <213> Homo sapiens

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 Arg Leu Arg Arg Pro Gly Gly Ser Tyr Pro Ala Ala Ala Ala Lys
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 Val Tyr Ser Leu Phe Arg Glu Gln Asp Ala Pro Val Ala Gly Leu Gln
 35 40 45
 Pro Val Glu Arg Ala Gln Pro Gly Trp Gly Ser Pro Arg Arg Pro Thr
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 Glu Ala Glu Ala Arg Arg Pro Ser Arg Ala Gln Gln Ser Arg Arg
 65 70 75

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 Val Tyr Ser Leu Phe Arg Glu Gln Asp Ala Pro Val Ala Gly Leu Gln
 35 40 45
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 50 55 60
 Glu Ala Glu Ala Arg Arg
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Val Tyr Ser Leu Phe Arg Glu Gln Asp Ala Pro Val Ala Gly Leu Gln
35 40 45
Pro Val Glu Arg Ala Gln Pro Gly Trp Gly Ser Pro Arg Arg Pro Thr
50 55 60
Glu Ala Glu Ala Arg Arg Pro Ser Arg Ala Gln Gln Ser Arg Arg
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Tyr Ser Leu Phe Arg Glu Gln Asp Ala Pro Val Ala Gly Leu Gln Pro
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Val Glu Arg Ala Gln Pro Gly Trp Gly Ser Pro Arg Arg
35 40 45

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Arg Pro Gly Gly Ser Tyr Pro Ala Ala Ala Ala Lys
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<211> 20
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1 5 10 15
Pro Val Glu Arg
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1 5 10

<210> 10

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<212> PRT

<213> Homo sapiens

<400> 10

Ala Gln Pro Gly Trp Gly Ser Pro Arg
1 5